

Immunohistochemistry of a Gross Cystic Disease Fluid Protein (GCDFP-15) of the Breast

A Marker of Apocrine Epithelium and Breast Carcinomas With Apocrine Features

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Gross cystic disease fluid is a pathologic secretion from breast composed of several glycoproteins, including a unique 15,000-dalton monomer protein, GCDFP-15. By the immunoperoxidase technique, GCDFP-15 was localized in the apocrine metaplastic epithelium lining breast cysts and in apocrine glands in the axilla, vulva, eyelid, and ear canal. In normal breast tissue, a few individual epithelial cells within lobules and small ducts were focally positive for GCDFP-15. Fourteen of 30 breast carcinomas stained positively for GCDFP-15. Of 16 carcinomas with apocrine features, 12 stained

positively. Benign and malignant lesions from other tissues, including lung, colon, ovary, endometrium, stomach, prostate, liver, esophagus, and kidney, revealed no immunoreactivity. The only cells of "non-apocrine" tissues that contained GCDFP-15 were serous cells of the submandibular salivary gland, submucosal glands of the bronchi, and accessory lacrimal glands. Phylogenetically, these tissues have biologic features in common with apocrine glands. This report is the first to characterize GCDFP-15 as a specific tissue marker of apocrine epithelium. (*Am J Pathol* 1983, 110:105-112)

GROSS CYSTIC DISEASE (GCD) of the breast is a premenopausal disorder in which gross cysts are the predominant pathologic lesion. Gross cysts appear to be formed from excessive apocrine cystic secretions.¹ The fluid found within these cysts is a unique secretion that only recently has been analyzed biochemically.²⁻⁷ Four major component proteins (gross cystic disease fluid proteins, or GCDFP) have consistently been identified in the cyst fluid: 1) GCDFP-70 (70,000 daltons), immunologically human albumin; 2) GCDFP-44, zinc alpha-2-glycoprotein; 3) GCDFP-24, which binds progesterone but not cortisol and accounts for over half of the protein content of GCD fluid; and 4) GCDFP-15, which has no normal plasma counterpart but, by immunodiffusion analysis, forms a line of identity with a normal component of saliva and milk.⁵

Cytosol analysis of normal tissue specimens from all major organs has demonstrated GCDFP-15 protein in apocrine epithelia. Detectable levels of GCDFP-15 are also present in the sublingual and

submaxillary salivary glands.⁸ Cytosols from breast carcinoma specimens also contain GCDFP-15 in a wide range of concentrations.⁸ The content is highest in more differentiated carcinomas and in those of intraductal cribriform and comedo types. Clinically, plasma levels of GCDFP-15 are useful in detecting and monitoring disease activity for approximately 40% of patients with metastatic breast carcinoma.⁴

The object of this study was to determine the specific tissue localization of GCDFP-15 in paraffin sections of normal tissues, nonneoplastic breast tissue with apocrine metaplasia, breast carcinomas with apocrine features, and a variety of other neoplastic tissue, with the use of an immunoperoxidase tech-

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nique. This report is the first to characterize GCDFP-15 as a specific tissue marker of apocrine cells and to define its immunohistochemical staining profile.

Materials and Methods

Paraffin blocks of normal tissues and of benign and malignant lesions of the breast and other organ systems were selected from the surgical pathology files (1975–1981) of Brigham and Women's Hospital. The specimens were initially fixed with either 10% neutral buffered formalin or Bouin's solution.

Identification of GCDFP-15 utilized a specific rabbit antiserum⁵ and the peroxidase–antiperoxidase immune complex method of Sternberger.⁹ Staining was performed at room temperature with reagents equilibrated to room temperature. Tissue sections were first deparaffinized, then rehydrated in xylene and graded alcohols. Endogenous peroxidase activity was blocked by a 30-minute incubation in 0.3% hydrogen peroxide,¹⁰ followed by a 2-minute incubation in 0.02% sodium borohydride.¹¹ After rinsing in phosphate-buffered saline (PBS) (0.15 M sodium chloride, 0.02 M sodium phosphate, pH 7.4), the tissues were incubated for 30 minutes with 10% normal swine serum (NSS) (Gibco, Grand Island, NY) in PBS. The slides were then drained of excess serum and wiped. The tissues were subsequently incubated for 2 hours with a 1:5000 dilution of the rabbit antiserum to GCDFP-15 diluted in 10% NSS. As a control, serial sections were incubated with a 1:1000 dilution of normal rabbit serum (NRS) diluted in 10% NSS. After rinsing with three changes of PBS for 10 minutes, a 1:40 dilution of swine anti-rabbit bridge antibody (Dako Corp, Santa Barbara, Calif) in 10% NSS was applied for 30 minutes. After rinsing, rabbit peroxidase–antiperoxidase (Dako Corporation) diluted 1:50 in 10% NSS was applied for 30 minutes. The sections were rinsed for 10 minutes in PBS; then a solution of 3,3'-diaminobenzidine tetrahydrochloride (DAB) (Sigma Chemical Company, St. Louis, Mo; 6 mg DAB in 10 ml PBS with 75 μ l hydrogen peroxide) was applied for a 3-minute reaction time. The sections were then washed in water and counterstained with hematoxylin, dehydrated in graded alcohols and xylene, and mounted with Permount.

A section of axillary skin (known positive) was included as a positive control in each experiment. Primary antisera absorbed against GCDFP-15 was also substituted for the immune serum on serial sections, as a negatively staining control for specificity of the antibody staining reaction. The antiserum was absorbed by the addition of 200 μ g of purified GCDFP-

15 to 5 ml antiserum diluted 1:5000 and incubated for at least 2 hours at room temperature.

Results

Immunoperoxidase studies of normal tissues stained for GCDFP-15 showed GCDFP-15 localization in the apocrine epithelium of several tissues: axilla, perineum, modified sweat glands (of Moll) of the eyelid, and the ceruminous glands of the external auditory canal (Table 1). Essentially all the apocrine cells in these tissues showed both discrete supranuclear and diffuse cytoplasmic staining for GCDFP-15 (Figures 1 and 2).

In benign breast tissues (2 normal, 1 fibroadenoma, 4 gross cystic disease, 1 adenoma, 2 gynecomastia), GCDFP-15 was predominantly localized within the metaplastic apocrine epithelium lining breast cysts and in the fluid contained within these cysts (Figure 3). All cysts with apocrine epithelium stained for GCDFP-15, from the smallest microcysts to gross cysts. Staining was most intense in discrete supranuclear areas of the cells. In the large cysts where the apocrine epithelium was flattened, GCDFP-15 staining was still retained.

In normal breast tissue, GCDFP-15 staining was observed only in a few individual epithelial cells within lobules and small ducts. Positive staining for GCDFP-15, however, did occur in a fibroadenoma that had areas of apocrine metaplasia.

Fourteen of 30 breast carcinomas stained positively for GCDFP-15 (Table 2). Sixteen of the 30 carcinomas had apocrine features in hematoxylin-and-eosin-stained sections, by the following morphologic criteria: abundant acidophilic cytoplasm, cytoplasmic "snouts," and large cell size in the routine sections.⁸ Twelve of these 16 carcinomas (75%) stained positively for GCDFP-15 and exhibited predominantly a diffuse cytoplasmic staining pattern (Figures 4 and 5), with an increased staining intensity adjacent

Table 1—Localization of GCDFP-15 in Normal Non-Mammary Tissues

Tissue	Immunoreactive Cells
Axillary skin (4)*	Apocrine sweat glands
Vulva, perineum (3)	Apocrine glands
External auditory canal (1)	Apocrine glands
Eyelid (1)	Apocrine glands
	Serous cells of lacrimal gland
Salivary gland	
Submandibular (3)	Serous cells
Parotid (7)	Equivocal (3 cases)
Lung (10)	Serous cells of bronchial glands

* Number of cases evaluated in parenthesis.

to the nucleus in some neoplastic cells. Four of these 12 cases showed GCDFP-15 immunoreactivity in the majority of malignant cells. In the remaining 8 cases, there was positivity in 10–25% of the neoplastic cells.

Of the 12 cases of breast carcinoma classified as intraductal, or infiltrating with an intraductal component, all had apocrine features, and 10 stained positively for GCDFP-15. Of 12 breast carcinomas classified as infiltrating carcinomas, 4 had apocrine features, and 2 of these stained positively for GCDFP-15. However, 2 other cases of infiltrating breast carcinoma, which by histologic criteria did not have apocrine features, also stained positively for GCDFP-15. None of the cases of infiltrating lobular carcinoma or mucoid carcinoma exhibited apocrine features or stained positively for GCDFP-15.

Normal tissue and malignant tumors from bladder, colon, endometrium, esophagus, kidney, liver, lung, ovary, pancreas, prostate, skin, stomach, thyroid, and tongue did not show any immunoreactivity for GCDFP-15, with the exception of apocrine sweat

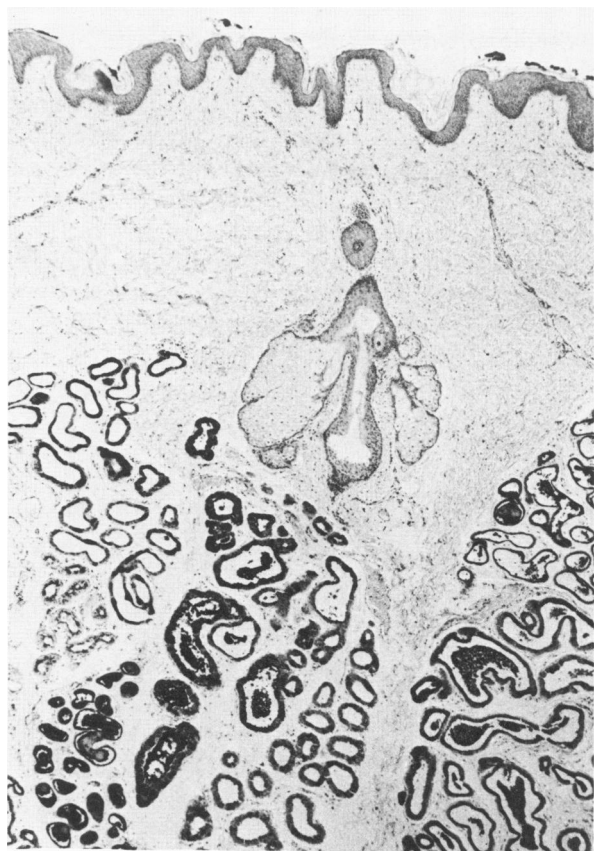


Figure 1—Skin, axilla. Immunoreactivity for GCDFP-15 is limited to apocrine glands and their luminal contents (*black*). Other skin appendages, epidermis, and dermal collagen are negative. (Hematoxylin counterstain, $\times 40$) (With a photographic reduction of 10%)



Figure 2—Skin, axilla. Higher magnification of apocrine glands demonstrates discrete supranuclear localization (*arrow*) of GCDFP-15 in many cells and focal diffuse cytoplasmic staining. (Hematoxylin counterstain, $\times 680$)

glands in the skin and the bronchial submucosal glands (Tables 3 and 4).

The only cells of “nonapocrine” tissues that contained GCDFP-15 were serous cells of the submandibular salivary gland, submucosal glands of the bronchi, and accessory lacrimal glands (Table 1; Figure 6). Sections of parotid glands, however, showed focal weak staining in the serous glands of 3 of 7 specimens.

Discussion

Our immunoperoxidase study demonstrates tissue localization of GCDFP-15 in normal apocrine cells of the body, benign breast epithelium that has undergone apocrine metaplasia, and breast carcinoma cells with apocrine features. Thus, morphologic features shared by a specific epithelium appear to be correlated with a biochemical product, GCDFP-15. This protein serves as an excellent tissue marker for apocrine epithelium.

The unique morphology of apocrine epithelium, with its granular acidophilic cytoplasm, has been studied by anatomists for over a century. In 1876 Krause¹² noted that the “sweat glands” in skin from the axilla, eyelid, and perianal area were larger than those in other regions of the body. Extending these

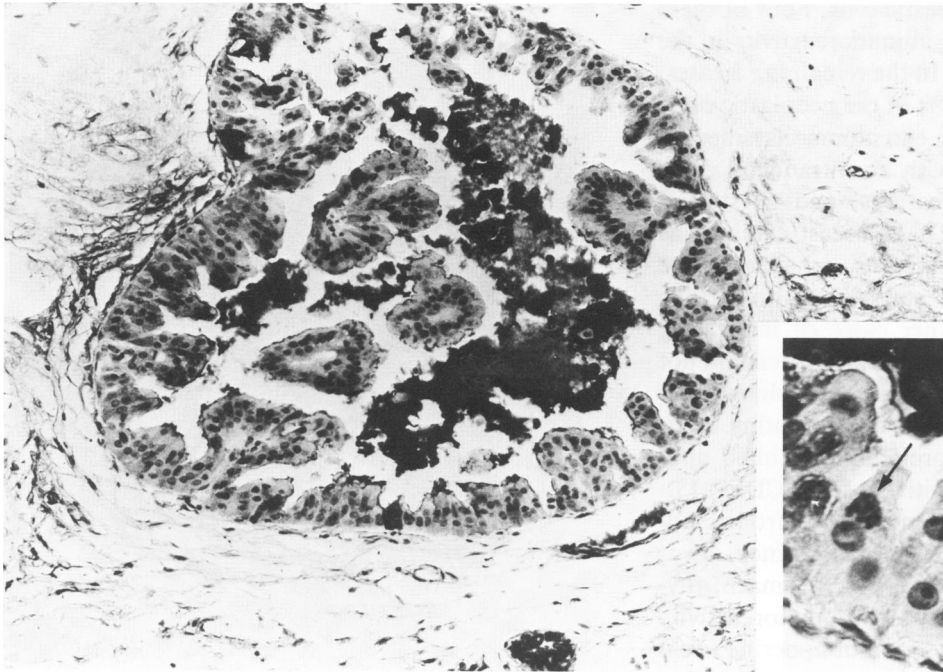


Figure 3—Breast, gross cystic disease. Immunoperoxidase staining of apocrine microcyst reveals strong positive staining of cyst fluid within the lumen for GCDFP-15. **Inset**—Localized supranuclear immunoreactivity (arrow) in epithelial cells lining the cyst. (Hematoxylin counterstain, $\times 140$; inset, $\times 750$)

observations, Schiefferdecker classified the cutaneous glands into holocrine (sebaceous) and merocrine.¹³ The latter he divided into “apocrine” and eccrine glands. Schiefferdecker described a mechanism of secretion of apocrine cells by decapitation or sheering off of the apical projections. This mechanism has recently been confirmed by ultrastructural study.¹⁴ Since the apocrine glands were a special type of gland under the influence of sex hormones, they have been considered as accessory sex glands or odoriferous glands.^{15,16}

Apocrine glands are found in the axilla, perineum, perianal region, the ceruminous glands, tarsal glands of the eyelid (Moll’s glands), and, lastly, the mammary glands.^{12,15,16} Benda¹⁷ postulated that the breast developed from a modified sweat gland. Krompecher¹⁸ recognized that the apocrine epithelium associated with breast cysts resembled the epithelium of the apocrine glands. A variety of hypotheses have been offered to explain the presence of apocrine epithelium in the breast: 1) Apocrine epithelium is a metaplasia or incomplete differentiation of normal breast epithelium into a less developed type.¹⁹ 2) Apocrine epithelium (“sweat glands”) is normally present in the breast.²⁰ 3) Apocrine epithelium is the terminal phase of breast epithelial degeneration, and the cells only simulate true apocrine epithelium and are not functionally related.²¹

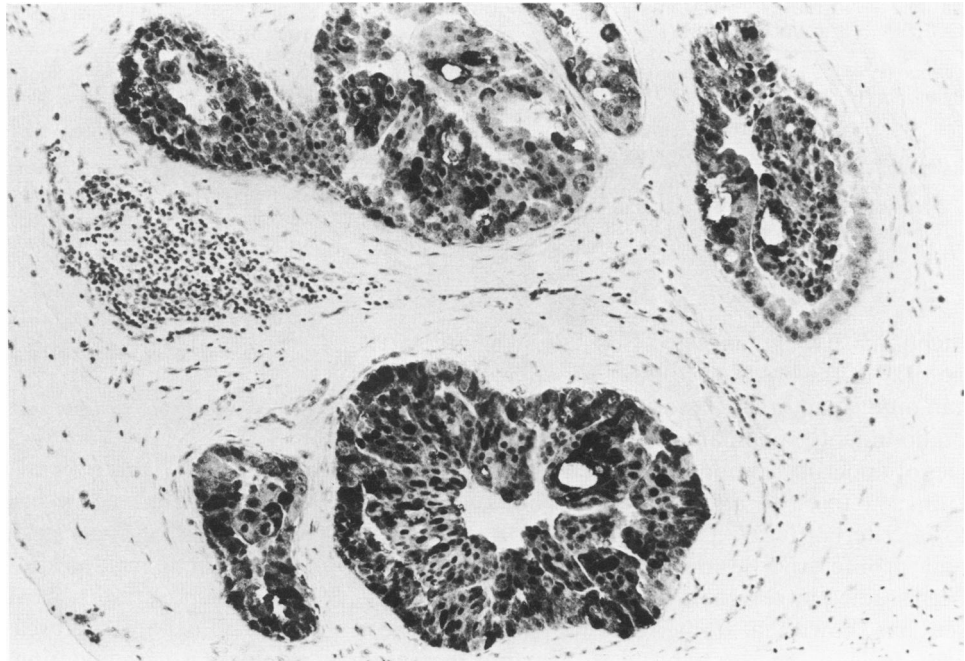
Studies of wholemount sections of breast have not demonstrated apocrine epithelium as a normal component of the breast duct system.²² Histochemical and ultrastructural studies,^{23–26} however, have shown characteristics of apocrine glands of the axilla to be similar to those of apocrine epithelium of the breast, supporting the hypothesis that it is a metaplastic phenomenon. Higginson and McDonald²⁷ compared the apocrine epithelium of the axilla with that of non-neoplastic breast tissue and apocrine carcinoma of the breast. They concluded in their study that “there is little doubt in our minds that the pale epithelium

Table 2—Breast Carcinomas Stained for GCDFP-15

Type of carcinoma	Number of cases	Cases with apocrine features	Number stained positively
Infiltrating	12	4	4*
Infiltrating and intraductal	4	4	3
Intraductal	8	8	7
Infiltration lobular (small-cell)	4	0	0
Mucoid	1	0	0
Cystosarcoma phyllodes	1	0	0
Total	30	16	14

* Two cases with apocrine features.

Figure 4 — Breast, adenocarcinoma. Most tumor cells of this intraductal carcinoma with papillary and apocrine features exhibit cytoplasmic staining for GCDFP-15. (Hematoxylin counterstain, $\times 170$)



found in breasts is actually apocrine glandular tissue identical with axillary apocrine scent glands . . . Truly conclusive proof would rest on a cellular biochemical study, with identification of secretions from the tissue.” The presence of a unique, secretory protein, GCDFP-15, in normal apocrine epithelium and

in apocrine epithelium of the breast is strong evidence for the latter representing a metaplastic process and further implies a functional relationship of these cells.

Krompecher was the first to speculate that some breast carcinomas were derived from apocrine epi-

Figure 5 — Breast, infiltrating ductal carcinoma. Immunoperoxidase staining for GCDFP-15 reveals cytoplasmic localization in tumor cells. (Hematoxylin counterstain, $\times 400$)

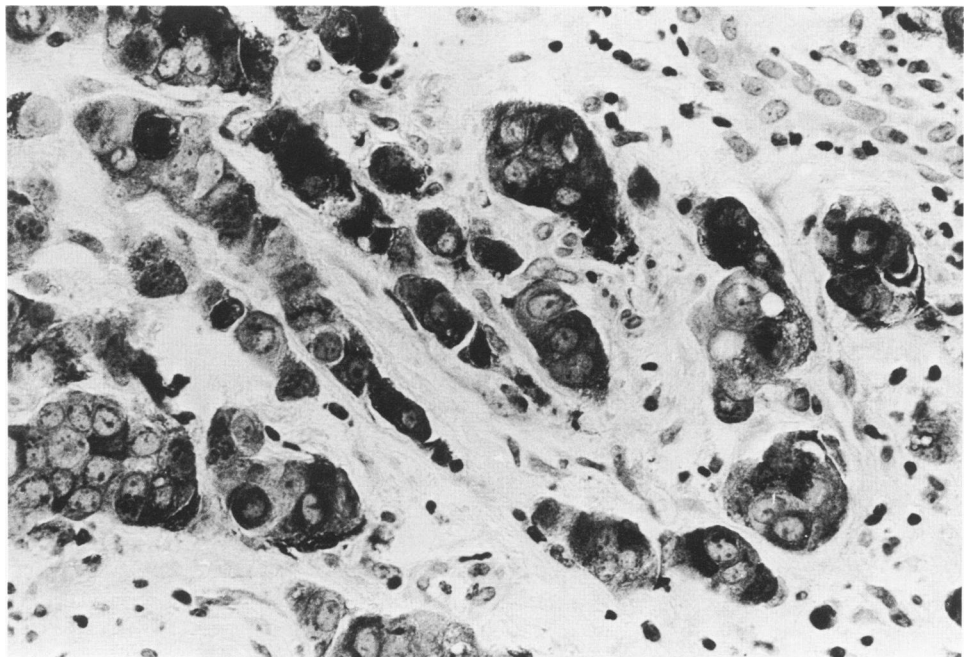


Table 3 — Normal Non-Mammary Tissues Without GCDFP-15 Immunoreactivity

Tongue (1)*	Pancreas (1)	Endometrium (2)
Epiglottis (1)	Colon (7)	Ovary (2)
Stomach (5)	Kidney (7)	Prostate (2)
Liver (2)	Bladder (1)	Thyroid (2)
Esophagus (1)		Lung† (10)

* Number of specimens evaluated in parenthesis.

† Immunoreactivity only in serous cells of bronchial glands.

thelium.²⁸ Ewing suggested that "sweat gland carcinomas" of the breast accounted for 25% of all breast carcinomas and that these apocrine carcinomas were predominantly papillary.²⁰ Other reported frequencies of apocrine carcinomas of the breast have ranged from 1% to 60%, reflecting the variability of histologic criteria.^{8,27,29-32} The relationship between cystic and proliferative lesions to breast carcinoma has been studied by several investigators.^{8,27,33-37} Haagen-⁸ has shown that women in whom gross cystic disease develops have a twofold to fourfold increased risk of breast carcinoma. He further noted a tenfold increase in the risk of carcinoma in patients with gross cystic disease with biopsy-proven microscopic apocrine metaplasia, as compared with those lacking this feature. The carcinomas that developed in the group with apocrine metaplasia were of apocrine

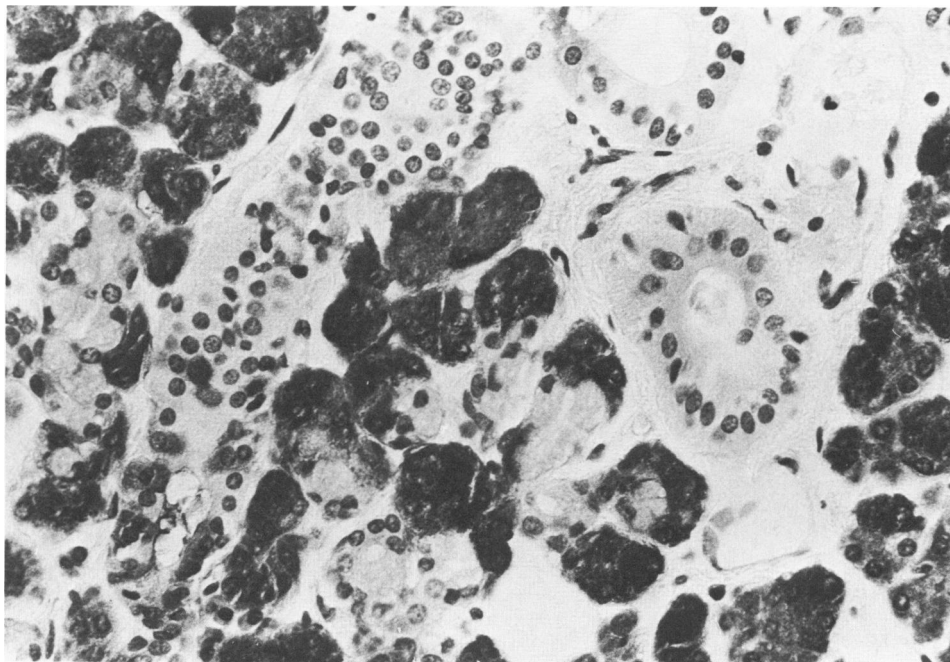
Table 4 — Non-Mammary Tumors Without GCDFP-15 Immunoreactivity

Adenocarcinoma	Squamous cell carcinoma
Esophagus (1)*	Lung (4)
Stomach (4)	Tongue (1)
Colon (5)	Basal cell carcinoma, skin (1)
Pancreas (2)	Transitional cell carcinoma, bladder (2)
Lung (3)	Cloacogenic carcinoma, rectum (2)
Kidney (5)	Hepatoma, liver (1)
Prostate (2)	Carcinoid, lung (2)
Endometrium (2)	Brenner tumor, ovary (1)
Ovary (2)	
Thyroid (1)	

* Numbers of cases evaluated in parenthesis.

type in 68%, and 65% of the carcinomas were intraductal. These studies support a hypothesis for the histogenesis of apocrine forms of breast carcinomas from benign metaplastic apocrine epithelium. Our study demonstrates localization of GCDFP-15 in apocrine metaplastic breast epithelium and in breast carcinomas with apocrine features but not in normal breast epithelium. These results provide biochemical evidence in favor of this histogenetic sequence.

The only cells of "nonapocrine" tissues containing GCDFP-15 were the serous cells of the submandibular gland, the lacrimal gland, and the bronchial mucosal glands. These tissues phylogenetically have fea-

**Figure 6 — Submandibular gland. GCDFP-15 immunoreactivity is distinctly localized to serous cells. Mucus cells and ductal epithelium are clearly negative. (Hematoxylin counterstain, ×340)**

tures in common with apocrine glands. Weak staining was also present in serous cells of some parotid glands. Parotid glands, however, may have a different embryologic origin than submandibular salivary glands.³⁸ It is of interest that in some species of animals the salivary glands function in the production of pheromones in a fashion similar to that of the apocrine glands.^{39,40} The salivary and lacrimal gland secretions are used by some species for marking territories and for inducing sexual stimulation. Androgens have been shown to be trophic to salivary glands also in a fashion similar to that of the apocrine glands.^{41,42}

Our immunohistochemical study suggests that GCDFP-15 production is a common embryologic and phylogenetic property of apocrine epithelium and that GCDFP-15 is a more sensitive indicator of apocrine epithelium than morphology alone. Immunoperoxidase staining for GCDFP-15 detects this biochemical characteristic of apocrine cells even when classic morphologic features are not apparent. The discrete paranuclear localization of GCDFP-15 probably corresponds to the Golgi complex region, which is well developed in breast apocrine metaplastic cells.²³

Our immunoperoxidase study for cellular localization of GCDFP-15 has identified a subgroup of breast carcinomas with apocrine features. The significance of this finding must be further assessed by correlation with other parameters of biologic activity such as estrogen and progesterone receptors, pattern of metastatic spread, and clinical response to therapy. This protein may represent a useful marker to suggest a possible breast origin for metastatic poorly differentiated neoplasms with unknown primary sites. Tissue localization of GCDFP-15 may also assist in defining or corroborating a possible apocrine derivation for skin tumors and in identifying subgroups of salivary gland neoplasms, particularly of submandibular origin.

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